

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

6544771

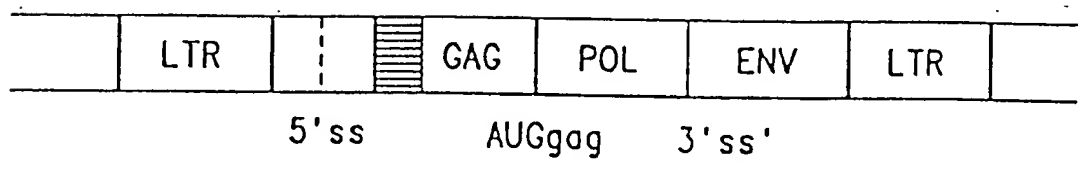


FIG. 1

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FIG. 2A pLJ

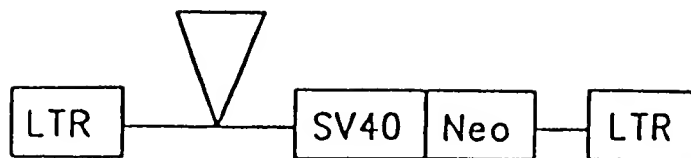


FIG. 2B pEm

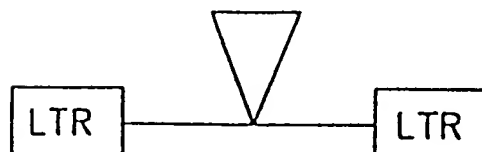


FIG. 2C MFG

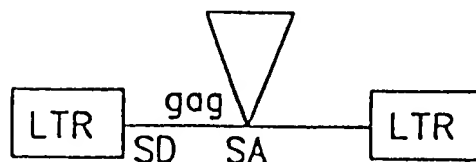
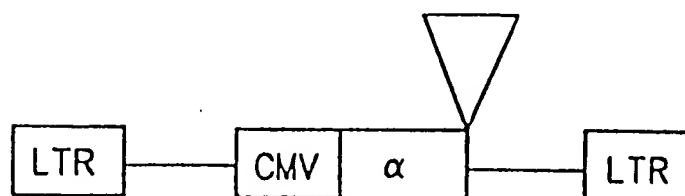


FIG. 2D α SGC



O.G. FIG.		
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		TS/MAN

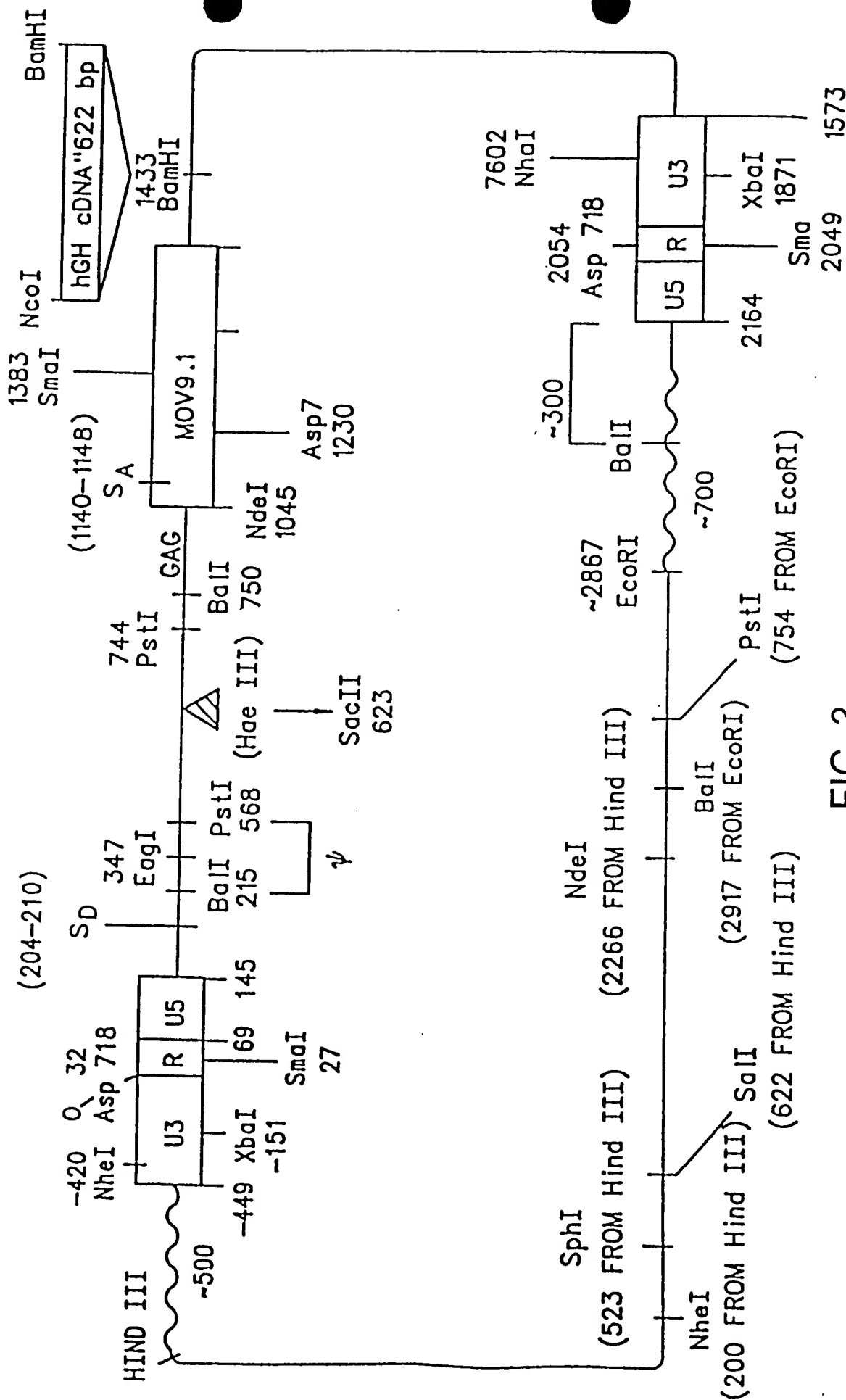
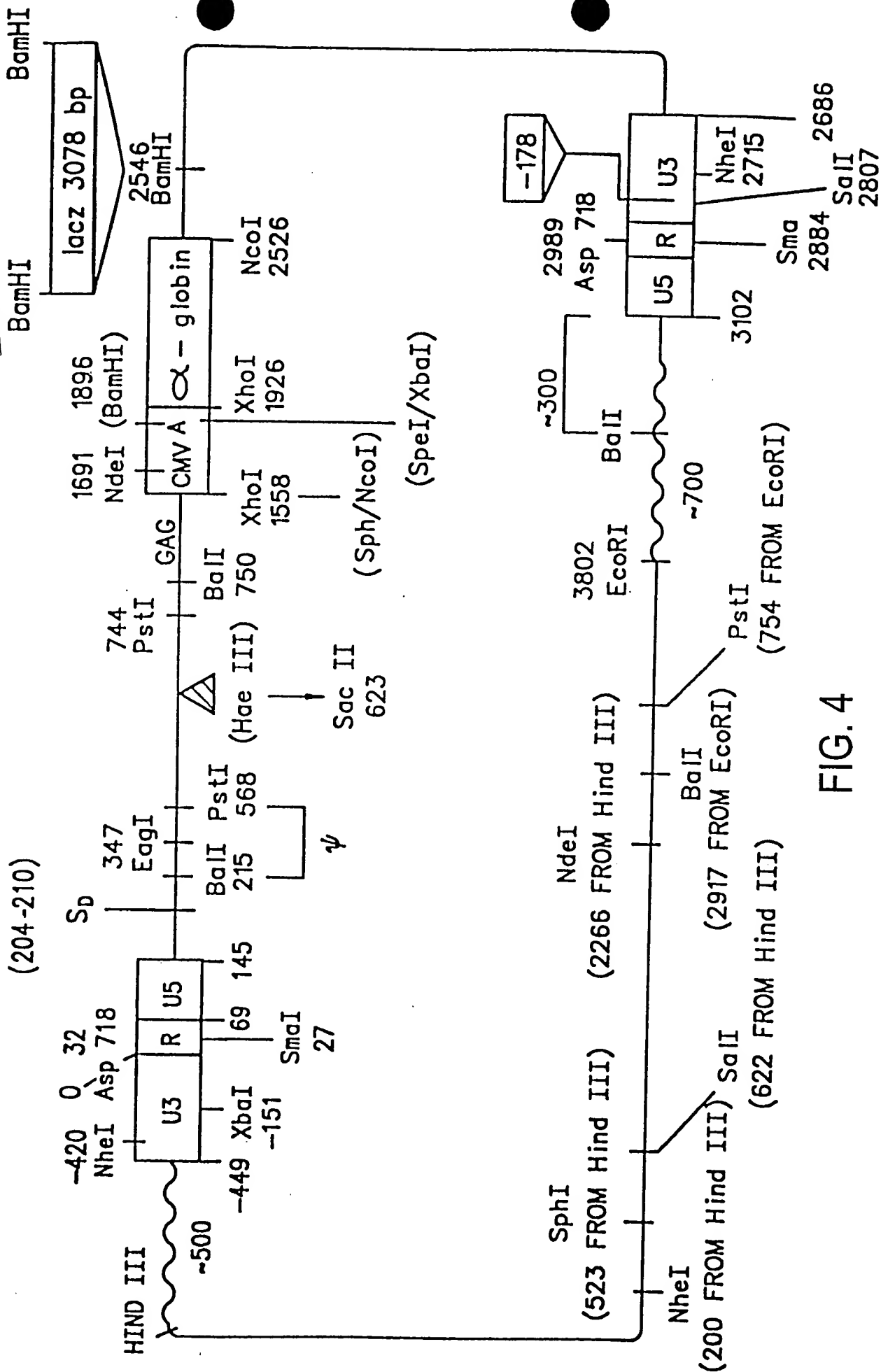


FIG. 3

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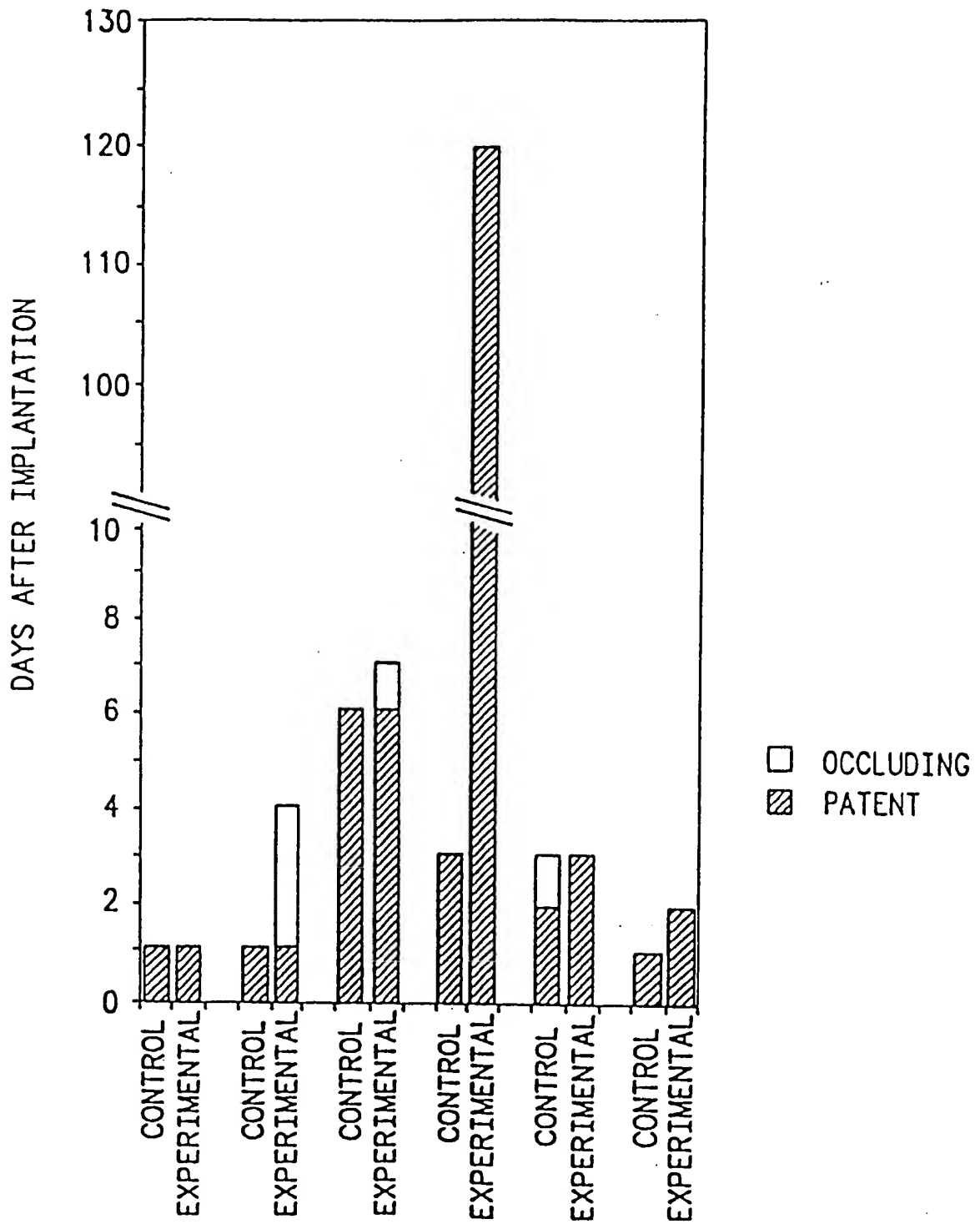


FIG. 5

O.G. FIG	CLASS	SUBCLASS
	PROVED BY	DATE

FIG. 6A

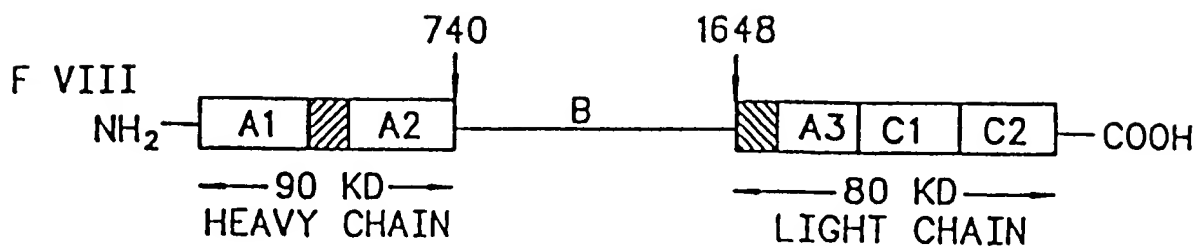


FIG. 6B

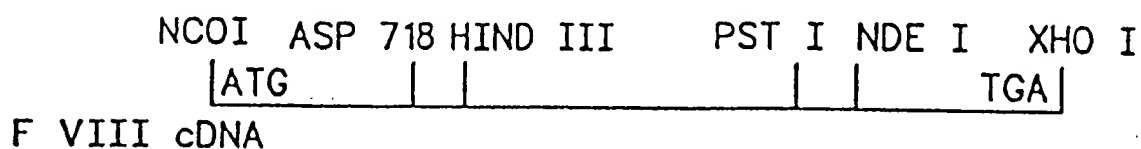


FIG. 6C

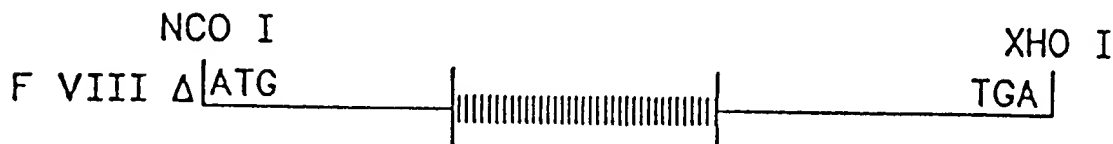
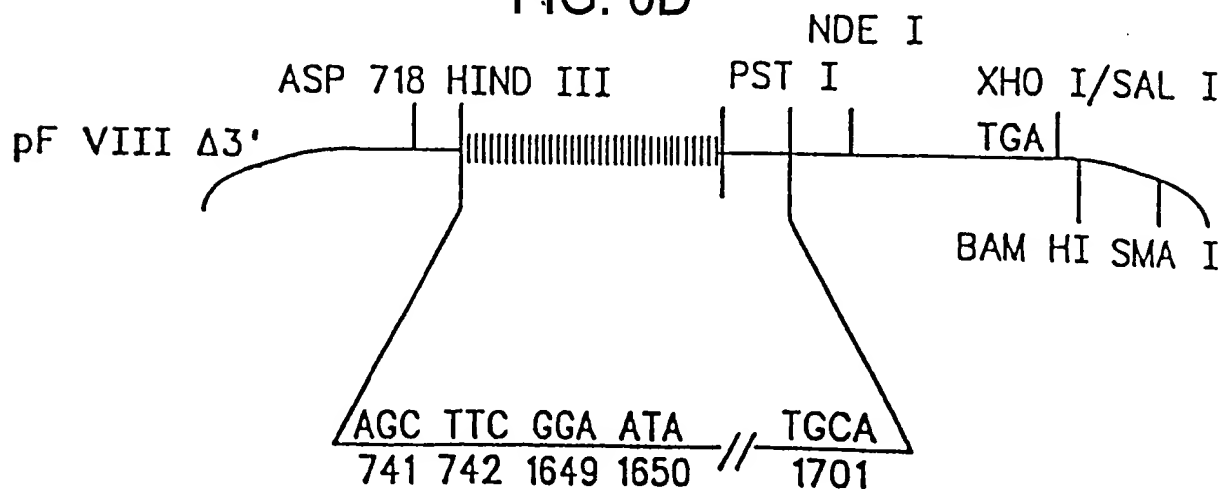


FIG. 6D



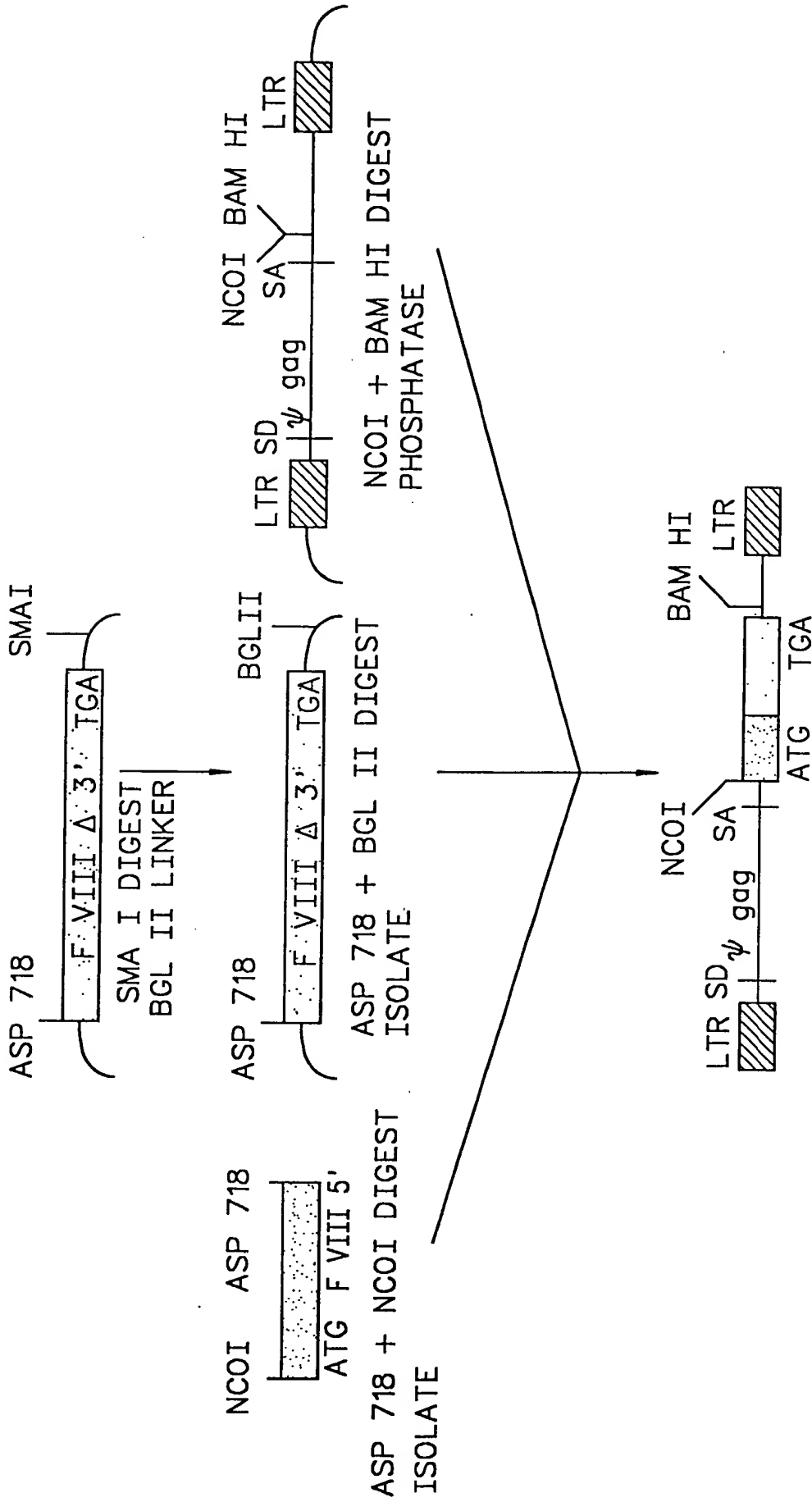
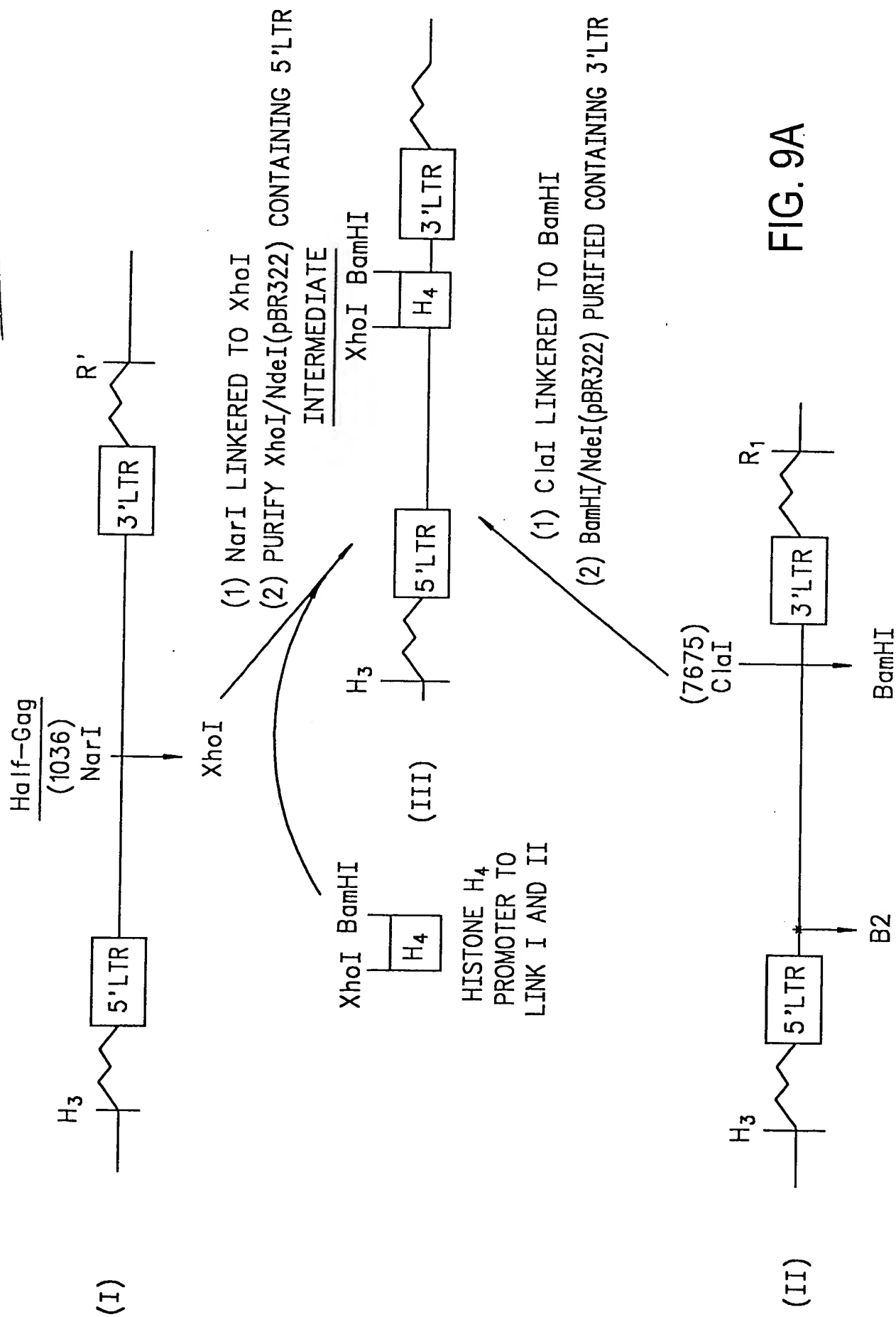


FIG. 7

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	CLASS	SUBCLASS



FIG. 8



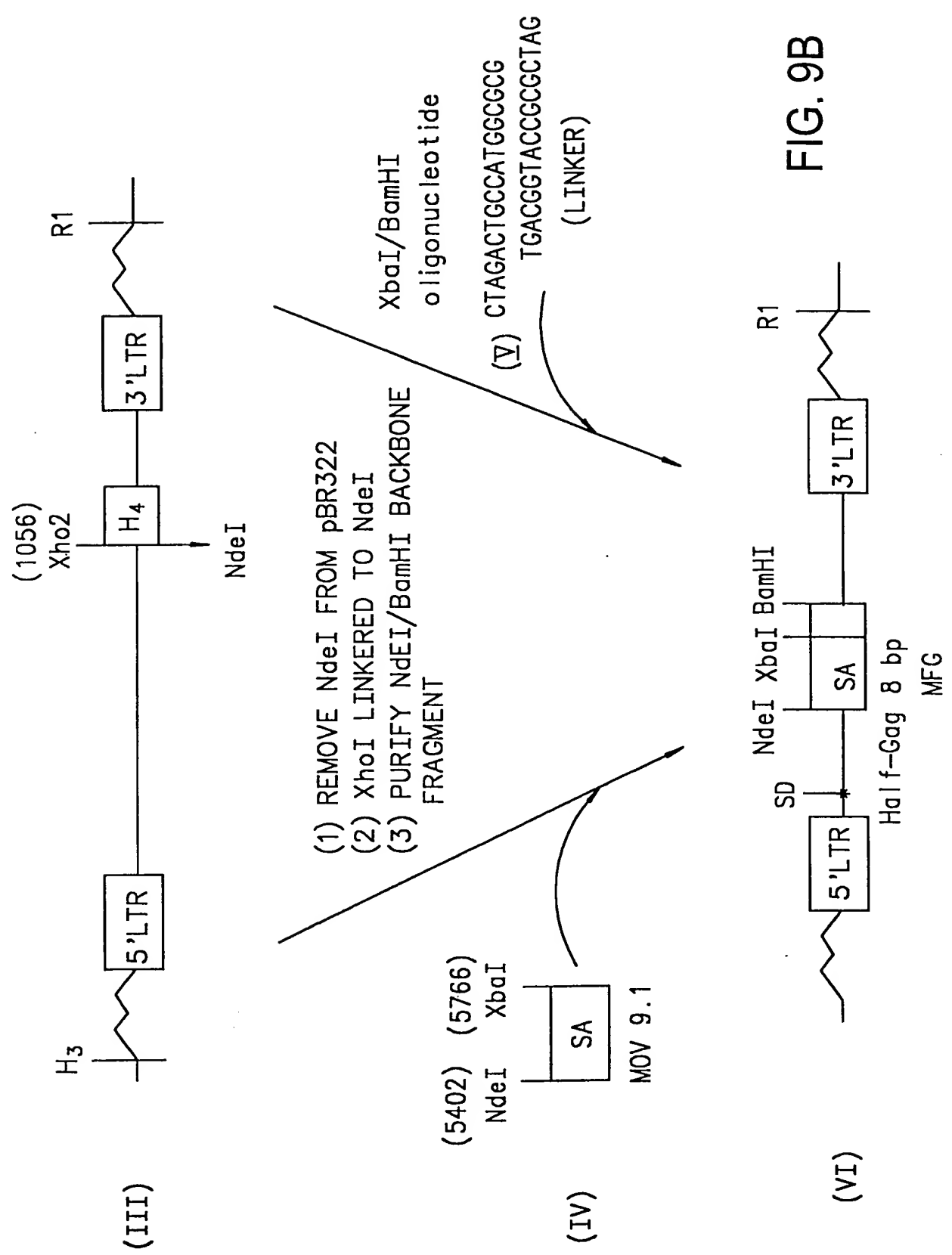
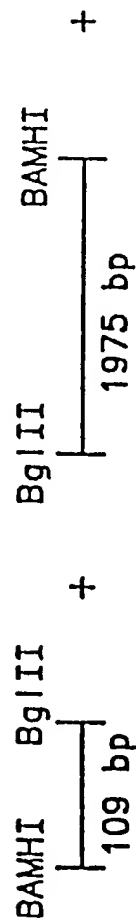
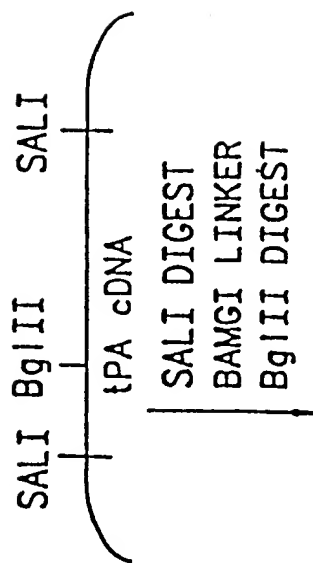


FIG. 9B



tPA oligo

```

5' C A T G G A T G C A A T G A A G A G A G
   NCOI 3' C T A C G T T A C T T C T C
        10
        20
        30
        40
        50
        60
        70
        80
        90
        100
G G C T C T G C T G T G T G C T G C T G
C C G A G A C G A C A C A C G A C G A C
C T G T G T G G A G C A G T C T T C G T
G A C A C A C C T C G T C A G A G C A
T T C G C C C A G C C A G G A A A T C C
A A G C G G G T C G G T C C C T T A G G
A T G C C C C G A T T C A G A A G A G A
T A C G G G G C T A A G T C T T C C C T
G C C A
C G G T C T A G

```

BglIII

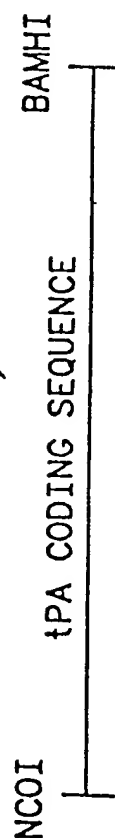
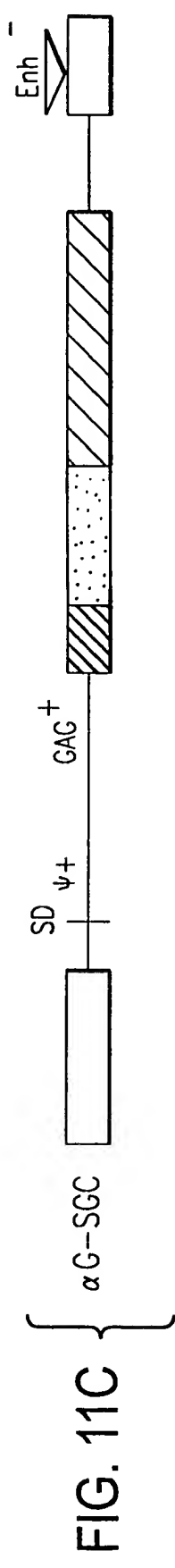
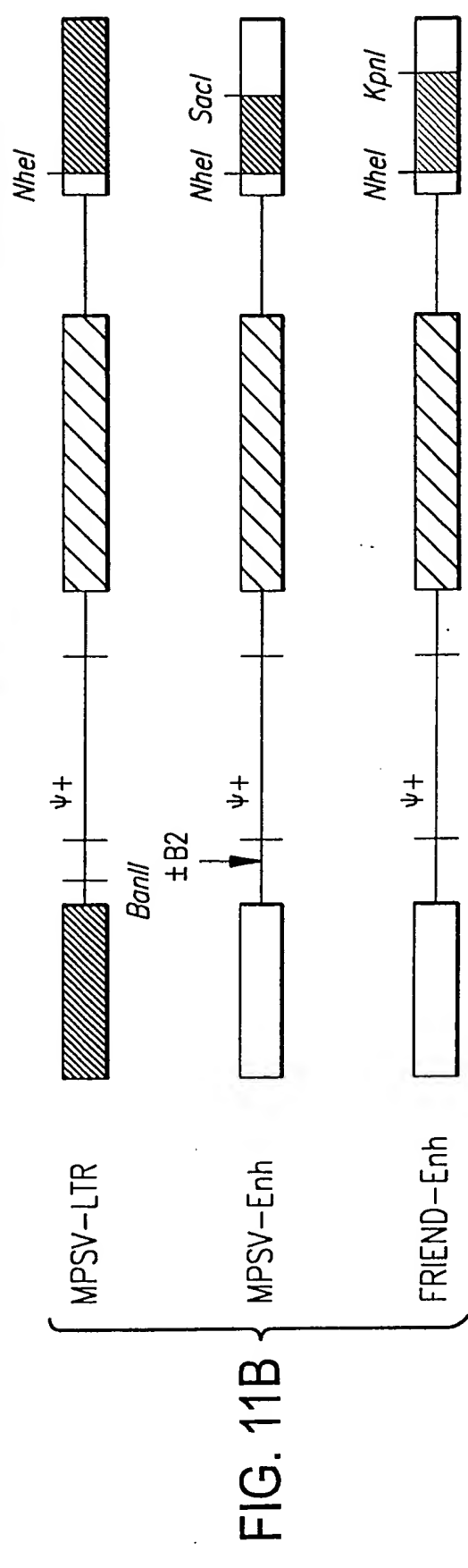
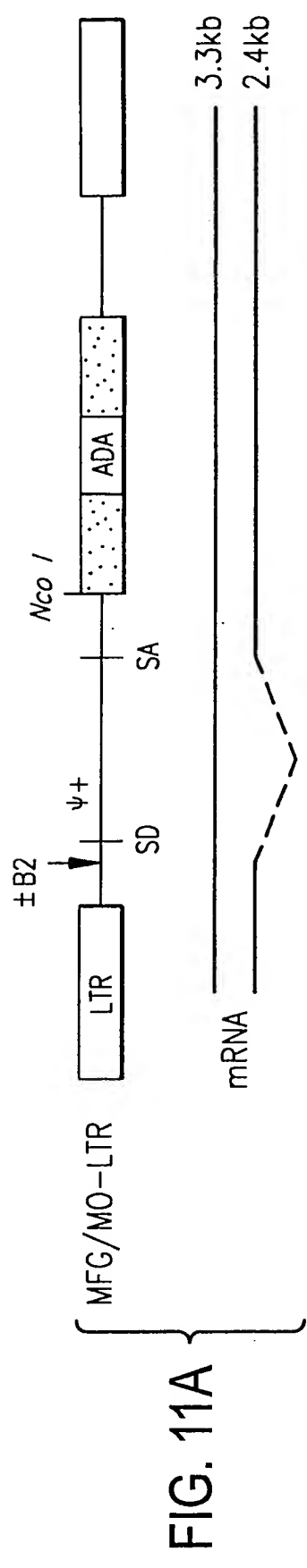


FIG. 10



	3×10^6																n_1	n_2	$n_1 + n_2/N$			
	5×10^5																$r \geq 1$	$1 < r \leq 1/4$				
Mo-LTR 191 DAYS	33	34	35	36	37	38	39	40	41	42	44	45	46	47	48	14	1	15/15				
hADA \blacktriangle mADA \blacktriangle	4.5×10^6																					
Mo-LTR/B2 191 DAYS	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	14	1	15/15				
	1.5×10^6																					
MPSV-Enh 206 DAYS	90	91	92	93	94	95	96	97	98	99	100	101	102	103	104	13	1	14/15				
	2×10^6																					
MPSV-Enh/B2 207 DAYS	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	9	6	15/15				
	10^6																					
Fr-Enh 184 DAYS	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	11	2	13/15				
	1.8×10^6																					
MPSV-LTR 177 DAYS	126	127	128	129	130	131	132	133	134	135	136	137	138	141	142	144	145	146	16	2	18/18	
	10^6																					
α G-SGC 170 DAYS	108	109	110	111	112	113	114	115	116	117	118	119	120	121	123	124	125	CONTROL		0	4	4/17
																	77	17	94/110			

FIC 10A

FIG. 12A

FIG. 12B

	DAYS AFTER BMT				hADA				mADA				RATIO hADA/mADA			
	191	402	191	402	191	402	191	402	191	402	191	402	191	402	191	402
Mo-LTR 80% (n=3)	#39	0.9	1.0	#40	1.4	0.7	#41	1.0	0.8							
Mo-LTR/B2	191	424	191	424	191	424	191	424	191	424	191	424	191	424	191	424
82% (n=4)	#54	1.8	1.5	#56	1.8	1.1	#57	1.8	2	#58	1.7	1.2				
Fr-Enh	184	430	184	430	184	430	184	430	184	430	184	430	184	430	184	430
80% (n=4)	#15	1.6	1.3	#16	1.4	0.8	#17	1.2	1.6	#18	1.2	0.6				
MPSV-LTR	177	361	177	361	177	361	177	361	177	361	177	361	177	361	177	361
77% (n=4)	#126	2.2	1.7	#127	2.2	2.0	#129	1.8	ND	#141	ND	ND	#142	2.5	1.5	#144
αG-SGC	170	420	170	420	170	420	170	420	170	399	170	420	170	420	170	420
59% (n=3)	#117	0.6	0.4	#121	0.05	0.03	#123	ND	ND	#118	0.8	0.4	CONTROL - NO VECTOR			

mADA

FIG. 12C

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- BM
- SPLEEN
- MAC
- T LYMPH
- B LYMPH

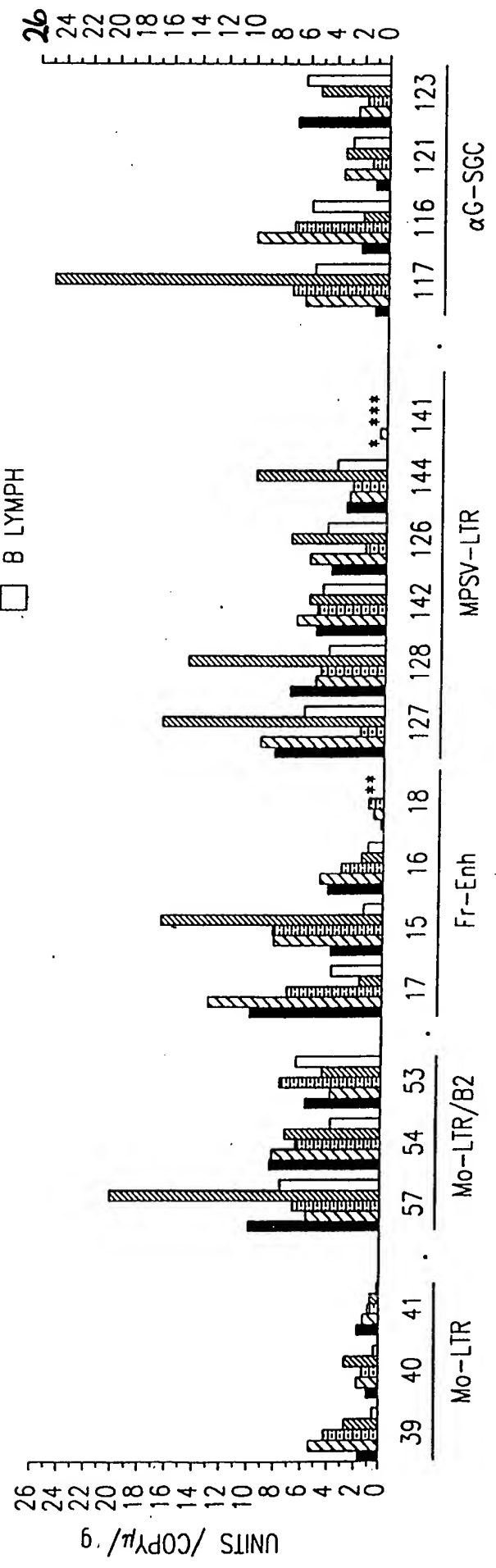


FIG. 13A

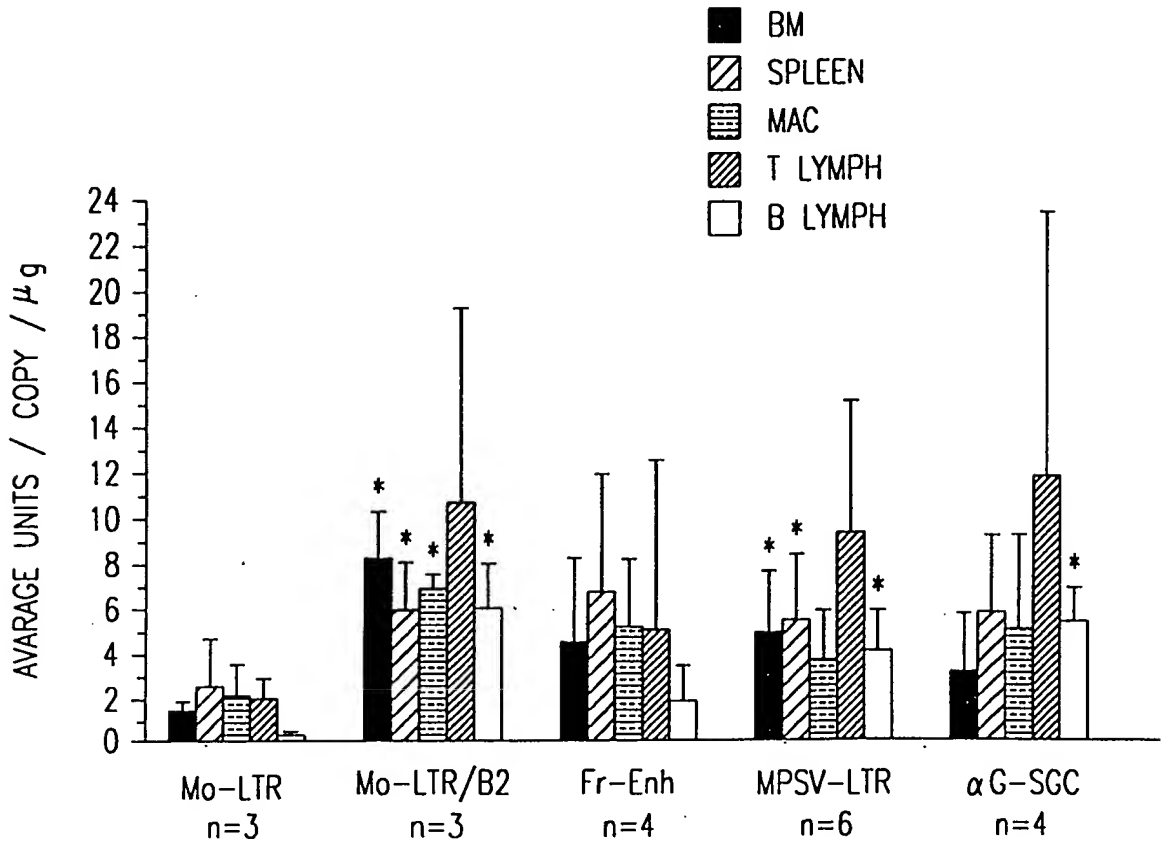


FIG. 13B

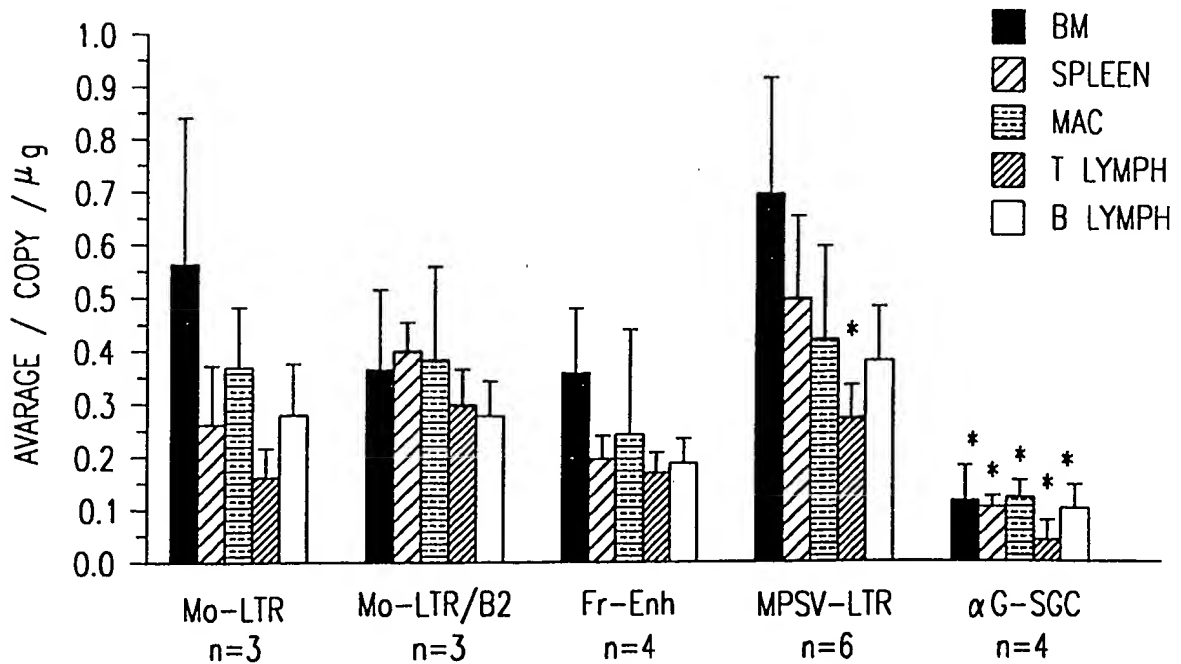


FIG. 13C

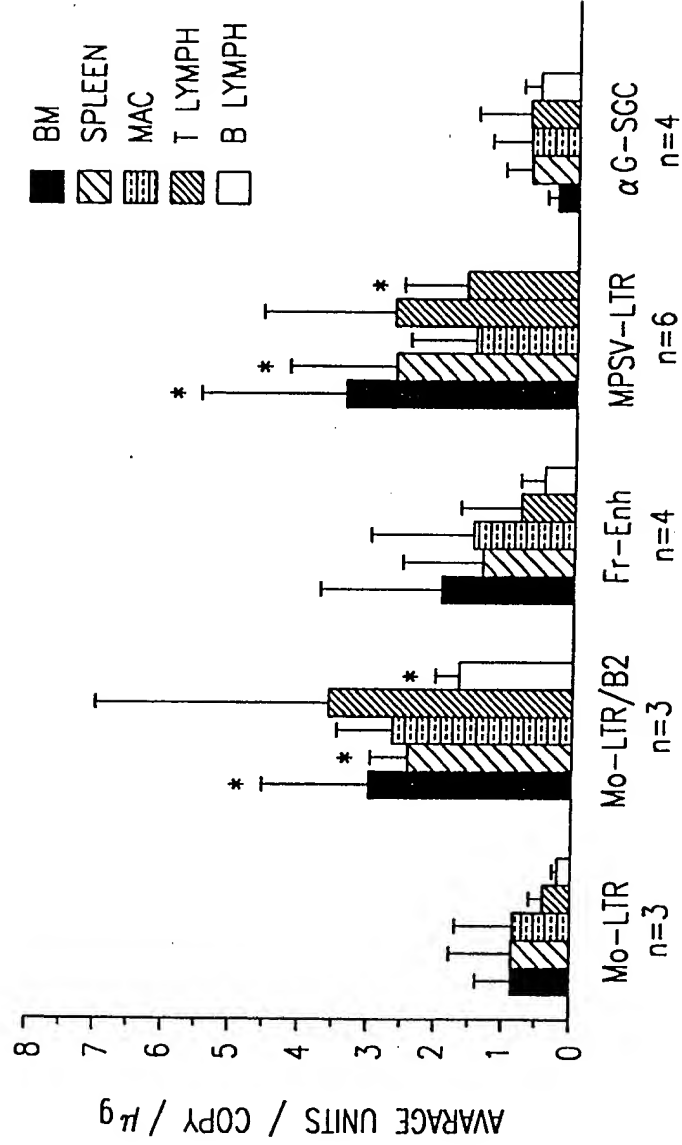


FIG. 13D

MoMuLV: GGTGGAACTGACGAGTTCGGAAACACCCGGCGCAACCTGGGAGACGTCCAGGGACTTCGGGGGCCGTTTTTIGIGCCCGACCT
MFG: XXXXXXXXXX
MFG-S: XXXXXXXXXX

MoMuLV: GAGTCCAAAAATCCCGATCGTTTTGGACICTTTGGTGCACCCCTTAGAGGAGGATAIGTGGTCTGGTAGGAGACGAGAACC
MFG: XX
MFG-S: XXXXXXXTXXX

MoMuLV: TAAACAGTCCCGCCCTCGTCTGAATTTTGGTTTGGGACCGAAGCCGGCGCGGTCTTGTCIGCTGCAGCATCGT
MFG: XX
MFG-S: XX

MoMuLV: ICIGTGTGICIGCTGACIGTGTTTICIGIATTTGTCIGAGAATAIGGG-----CCAGACTGTACCACTCCCT
MFG: XX
MFG-S: XX

FIG. 14

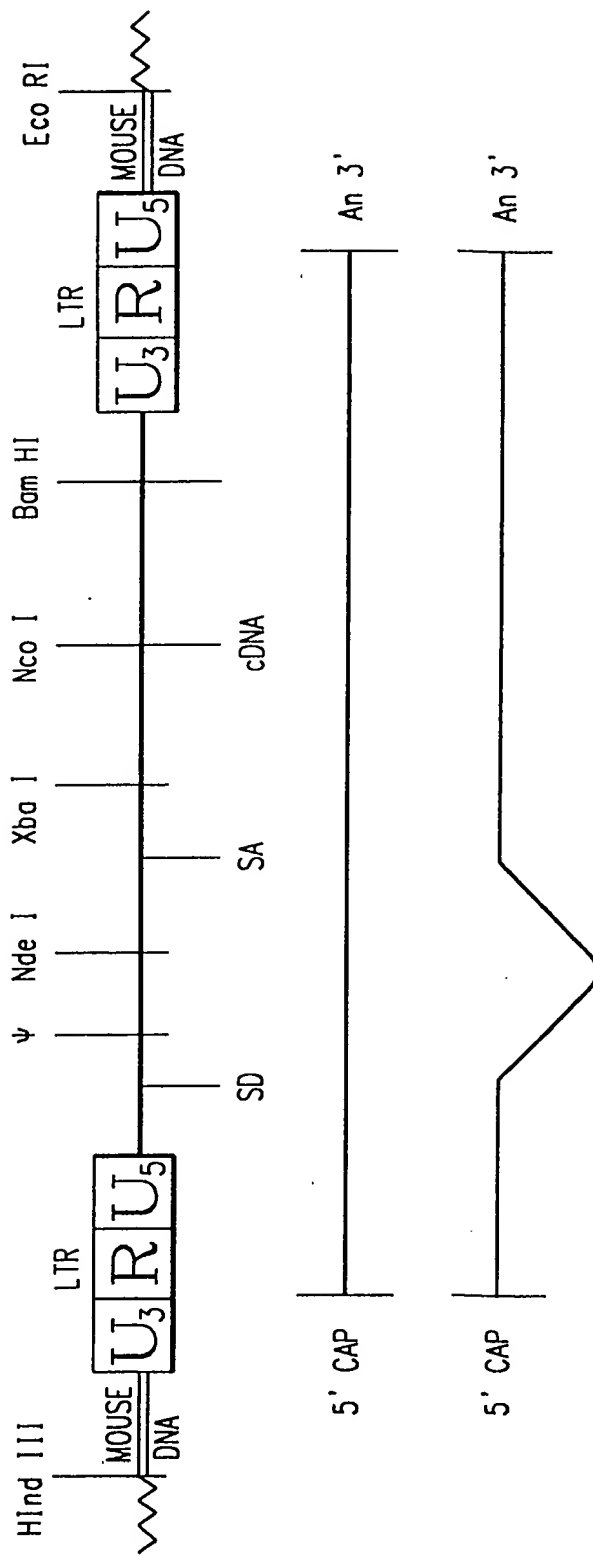


FIG. 15

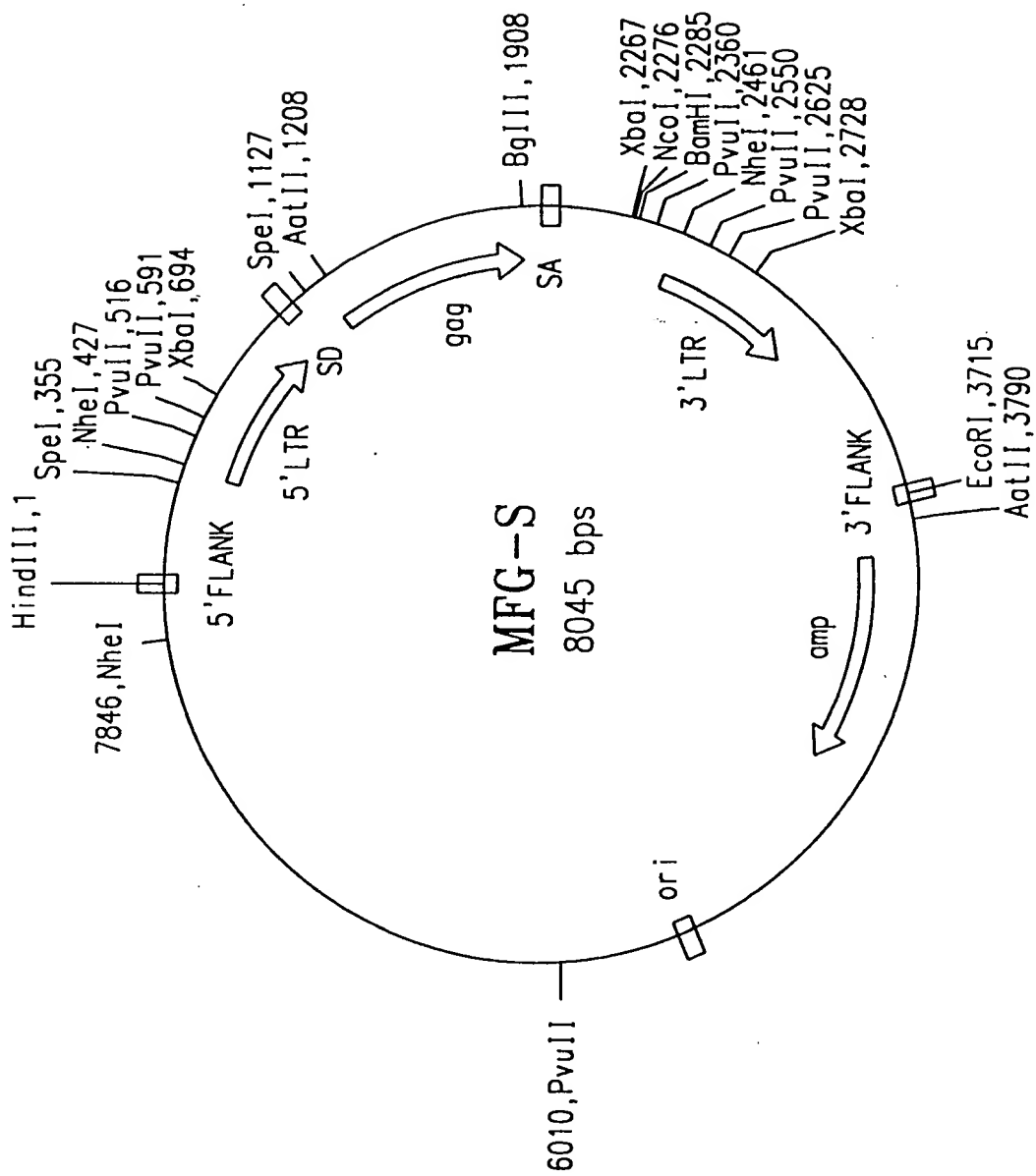


FIG. 16

1 AAGCTTTGCT CTTAGGAGTT TCCTAATACA TCCCAAACCTC AAATATATAA AGCATTTGAC
TTCGAAACGA GAATCCTCAA AGGATTATGT AGGGTTTGAG TTTATATATT TCGTAAACTG

61 TTGTTCTATG CCCTAGGGGG CGGGGGGAAG CTAAGCCAGC TTTTTTTAAC ATTTAAAATG
AACAAGATAC GGGATCCCCC GCCCCCTTC GATTCCGTG AAAAAAATTG TAAATTTTAC

121 TTAATTCCAT TTAAATGCA CAGATGTTTT TATTTCATAA GGGTTTCAAT GTGCATGAAT
AATTAAGGTA AAATTTACGT GTCTACAAAA ATAAAGTATT CCCAAAGTTA CACGTACTTA

181 GCTGCAATAT TCCTGTTACC AAAGCTAGTA TAAATAAAAA TAGATAAACG TGGAAATTAC
CGACGTTATA AGGACAATGG TTTGATCAT ATTTATTTTT ATCTATTTGC ACCTTTAATG

241 TTAGAGTTTC TGTCAATTAAC GTTTCCTTCC TCAGTTGACA ACATAAATGC GCTGCTGAGC
AATCTCAAAG ACAGTAATTG CAAAGGAAGG AGTCAACTGT TGTATTTACG CGACGACTCG

301 AAGCCAGTTT GCATCTGTCA GGATCAATTT CCCATTATGC CAGTCATATT AATTACTAGT
TTCGGTCAAA CGTAGACAGT CCTAGTTAAA GGGTAATACG GTCAGTATAA TTAATGATCA

361 CAATTAGTTG ATTTTTATTT TTGACATATA CATGTGAATG AAAGACCCCA CCTGTAGGTT
GTAAATCAAC TAAAAATAAA AACTGTATAT GTACACTTAC TTTCTGGGGT GGACATCCAA

421 TGGCAAGCTA GCTTAAGTAA CGCCATTTTG CAAGGCATGG AAAAATACAT AACTGAGAAT
ACCGTTCCAT CGAATTCATT GCGGTAAAC GTTCCGTACC TTTTATGTA TTGACTCTTA

481 AGAAAAGTTC AGATCAAGGT CAGGAACAGA TGAACAGCT GAATATGGGC CAAACAGGAT
TCTTTTCAAG TCTAGTTCCA GTCCTTGCT ACCTTGTCGA CTTATACCCG GTTGTGCTTA

541 ATCTGTGGTA AGCAGTTCCT GCCCCGGCTC AGGGCCAAGA ACAGATGGAA CAGCTGAATA
TAGACACCAT TCGTCAAGGA CGGGGCCGAG TCCCGTTCT TGTCTACCTT GTCGACTTAT

601 TGGGCCAAAC AGGATATCTG TGTAAGCAG TTCCTGCCCC GGCTCAGGGC CAAGAACAGA
ACCCGGTTTG TCCTATAGAC ACCATTGCTC AAGGACGGGG CCGAGTCCCG GTTCTTGTCT

661 TGGTCCCCAG ATGCGGTCCA GCCCTCAGCA GTTCTAGAG AACCATCAGA TGTTCACAGG
ACCAGGGGTC TACGCCAGGT CGGGAGTCGT CAAAGATCTC TTGGTAGTCT ACAAAGGTCC

721 GTGCCCCAAG GACCTGAAAT GACCCTGTGC CTTATTTGAA CTAACCAATC AGTTGCTTC
CACGGGGTTC CTGGACTTTA CTGGGACACG GAATAAACTT GATTGGTTAG TCAAGCGAAG

781 TCGCTTCTGT TCGCGCGCTT CTGCTCCCCG AGCTCAATAA AAGAGCCCAC AACCCCTCAC
AGCGAAGACA AGCGCGCGAA GACGAGGGGC TCGAGTTATT TTCTCGGGTG TTGGGGAGTG

FIG. 17A

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841 TCGGGGCGCC AGTCCTCCGA TTGACTGAGT CGCCCGGGTA CCCGTGTATC CAATAAACCC
 AGCCCCGCGG TCAGGAGGCT AACTGACTCA GCGGGCCCAT GGGCACATAG GTTATTTGGG
 901 TCTTGCAGTT GCATCCGACT TGTGGTCTCG CTGTTCTTGG GGAGGGTCTC CTCTGAGTGA
 AGAACGTCAA CGTAGGCTGA ACACCAGAGC GACAAGGAAC CCTCCCAGAG GAGACTCACT
 961 TTGACTACCC GTCAGCGGGG GTCTTTTCATT TGGGGGCTCG TCCGGGATCG GGAGACCCCT
 AACTGATGGG CAGTCGCCCC CAGAAAGTAA ACCCCCGAGC AGGCCCTAGC CCTCTGGGGA
 1021 GCCCAGGGAC CACCGACCCA CCACCGGGAG GTAAGCTGGC CAGCAACTTA TCTGTGTCTG
 CCGGTCCCTG GTGGCTGGGT GGTGGCCCTC CATTCGACCG GTCGTTGAAT AGACACAGAC
 1081 TCCGATTGTC TAGTGTCTAT GACTGATTTT ATGCGCCTGC GTCGGTACTA GTTAGCTAAC
 AGGCTAACAG ATCACAGATA CTGACTAAAA TACGCGGACG CAGCCATGAT CAATCGATTG
 1141 TAGCTCTGTA TCTGGCGGAC CCGTGGTGA ACTGACGAGT TCGGAACACC CGGCCGCAAC
 ATCGAGACAT AGACCGCCTG GGCACCACCT TGA CTGCTCA AGCCTTGTGG GCCGGCGTTG
 1201 CCTGGGAGAC GTCCAGGGA CTTCGGGGGC CGTTTTTGTG GCCCGACCTG AGTCCTAAAA
 GGACCCTCTG CAGGGTCCCT GAAGCCCCCG GCAAAAACAC CGGGCTGGAC TCAGGATTTT
 1261 TCCCGATCGT TTAGGACTCT TTGGTGCACC CCCCTTAGAG GAGGGATATG TGGTTCTGGT
 AGGGCTAGCA AATCCTGAGA AACCACGTGG GGGGAATCTC CTCCCTATAC ACCAAGACCA
 1321 AGGAGACGAG AACCTAAAAC AGTTCCCGCC TCCGTCTGAA TTTTGTCTTT CGGTTTGGGA
 TCCTCTGCTC TTGGATTTTG TCAAGGGCGG AGGCAGACTT AAAAACGAAA GCCAAACCCT
 1381 CCGAAGCCGC GCCGCGCGTC TTGTCTGCTG CAGCATCGTT CTGTGTTGTC TCTGTCTGAC
 GGCTTCGGCG CGGCGCGCAG AACAGACGAC GTCGTAGCAA GACACAACAG AGACAGACTG
 1441 TGTGTTTCTG TATTTGTCTG AAAATATGGG CCCGGGCTAG ACTGTTACCA CTCCCTTAAG
 ACACAAAGAC ATAAACAGAC TTTTATACCC GGGCCCGATC TGACAATGGT GAGGGAATTC
 1501 TTTGACCTTA GGTCAGTGA AAGATGTCCA GCGGATCGCT CACAACCAGT CGGTAGATGT
 AAAGTGAAT CCAGTGACCT TTCTACAGCT CGCCTAGCGA GTGTTGGTCA GCCATCTACA
 1561 CAAGAAGAGA CGTTGGGTTA CCTTCTGCTC TGCAGAATGG CCAACCTTTA ACGTCGGATG
 GTTCTTCTCT GCAACCCAAT GGAAGACGAG ACGTCTTACC GGTGGAAT TGCAGCCTAC

FIG. 17B

1621 GCCGCGAGAC GGCACCTTTA ACCGAGACCT CATCACCCAG GTTAAGATCA AGGTCTTTTC
CGGCGCTCTG CCGTGGAAAT TGGCTCTGGA GTAGTGGGTC CAATTCTAGT TCCAGAAAAG

1681 ACCTGGCCCG CATGGACACC CAGACCAGGT CCCCTACATC GTGACCTGGG AAGCCTTGGC
TGGACCGGGC GTACCTGTGG GTCTGGTCCA GGGGATGTAG CACTGGACCC TTCGGAACCG

1741 TTTTGACCCC CCTCCCTGGG TCAAGCCCTT TGTACACCTT AAGCCTCCGC CTCCTCTTCC
AAACTGGGG GGAGGGACCC AGTTCGGGAA ACATGTGGGA TTCGGAGGCG GAGGAGAAGG

1801 TCCATCCGCC CCGTCTCTCC CCCTTGAACC TCCTCGTTCC ACCCCGCCTC GATCCTCCCT
AGGTAGGCGG GGCAGAGAGG GGGAACTTGG AGGAGCAAGC TGGGGCGGAG CTAGGAGGGA

1861 TTATCCAGCC CTCACCTCTT CTCTAGGCGC CCCCATATGG CCATATGAGA TCTTATATGG
AATAGGTCCG GAGTGAGGAA GAGATCCGCG GGGGTATACC GGTATACTCT AGAATATACC

1921 GGCACCCCGG CCCCTTGTA ACTTCCCTGA CCCTGACATG ACAAGAGTTA CTAACAGCCC
CCGTGGGGGC GGGGAACATT TGAAGGACT GGGACTGTAC TGTTCTCAAT GATTGTGGG

1981 CTCTCTCCAA GCTCACTTAC AGGCTCTCTA CTTAGTCCAG CACGAAGTCT GGAGACCTCT
GAGAGAGGTT CGAGTGAATG TCCGAGAGAT GAATCAGGTC GTGCTTCAGA CCTCTGGAGA

2041 GGCGGCAGCC TACCAAGAAC AACTGGACCG ACCGGTGGTA CCTCACCTT ACCGAGTCGG
CCCCGTCGG ATGTTCTTG TTGACCTGGC TGGCCACCAT GGAGTGGAA TGGCTCAGCC

2101 CGACACAGTG TGGGTCCGCC GACACCAGAC TAAGAACCTA GAACCTCGCT GGAAAGGACC
GCTGTGTCAC ACCCAGGCGG CTGTGGTCTG ATTCTTGGAT CTTGGAGCGA CCTTTCCTGG

2161 TTACACAGTC CTGCTGACCA CCCCCACCGC CCTCAAAGTA GACGGCATCG CAGCTTGGAT
AATGTGTCAG GACGACTGGT GGGGGTGGCG GGAGTTTCAT CTGCCGTAGC GTCGAACCTA

2221 ACACGCCGCC CACGTGAAGG CTGCCGACCC CGGGGGTGGG CCATCCTCTA GACTGCCATG
TGTGGGGCGG GTGCACTTCC GACGGCTGGG GCCCCACCT GGTAGGAGAT CTGACGGTAC

2281 GCGCGGATCC GGATTAGTCC AATTTGTTAA AGACAGGATA TCAGTGGTCC AGGCTCTAGT
CGCGCCTAGG CCTAATCAGG TTAAACAATT TCTGTCCTAT AGTCACCAGG TCCGAGATCA

2341 TTTGACTCAA CAATATCACC AGCTGAAGCC TATAGAGTAC GAGCCATAGA TAAAATAAAA
AAACTGAGTT GTTATAGTGG TCGACTTCGG ATATCTCATG CTCGGTATCT ATTTTATTTT

FIG. 17C

2401 GATTTTATTT AGTCTCCAGA AAAAGGGGGG AATGAAAGAC CCCACCTGTA GGTITGGCAA
CTAAAATAAA TCAGAGGTCT TTTTCCCCC TTACTTTCTG GGGTGGACAT CCAAACCGTT

2461 GCTAGCTTAA GTAACCCCAT TTTGCAAGGC ATGGAAAAAT ACATAACTGA GAATAGAGAA
CGATCGAATT CATTGCGGTA AAACGTTCCG TACCTTTTAA TGTATTGACT CTTATCTCTT

2521 GTTCAGATCA AGGTCAGGAA CAGATGGAAC AGCTGAATAT GGGCCAAACA GGATATCTGT
CAAGTCTAGT TCCAGTCCTT GTCTACCTTG TCGACTTATA CCCGGTTTGT CCTATAGACA

2581 GGTAAGCAGT TCCTGCCCCG GCTCAGGGCC AAGAACAGAT GGAACAGCTG AATATGGGCC
CCATTGCTCA AGGACGGGGC CGAGTCCCGG TTCTTGTCTA CCTTGTGCGAC TTATACCCGG

2641 AAACAGGATA TCTGTGGTAA GCAGTTCCTG CCCC GGCTCA GGGCCAAGAA CAGATGGTCC
TTTGTCTAT AGACACCAT CGTCAAGGAC GGGGCCGAGT CCCGGTTCTT GTCTACCAGG

2701 CCAGATGCGG TCCAGCCCTC AGCAGTTTCT AGAGAACCAT CAGATGTTTC CAGGGTGCCC
GGTCTACGCC AGGTCGGGAG TCGTCAAAGA TCTCTTGGTA GTCTACAAAG GTCCACGGG

2761 CAAGGACCTG AAATGACCCT GTGCCTTATT TGAAC TAACC AATCAGTTCG CTTCTCGCTT
GTTCTGGAC TTTACTGGGA CACGAATAA ACTTGATTGG TTAGTCAAGC GAAGAGCGAA

2821 CTGTTGCGGC GCTTCTGCTC CCCGAGCTCA ATAAAAGAGC CCACAACCCC TACTCGGGG
GACAAGCGCG CGAAGACGAG GGGCTCGAGT TATTTTCTCG GGTGTTGGG AGTGAGCCCC

2881 CGCCAGTCCT CCGATTGACT GAGTCGCCCC GGTACCCGTG TATCCAATAA ACCCTCTTGC
GCGGTCAGGA GGCTAACTGA CTCAGCGGGC CCATGGGCAC ATAGGTTATT TGGGAGAACG

2941 AGTTGCATCC GACTTGTGGT CTCGCTGTTT CTTGGGAGGG TCTCCTCTGA GTGATTGACT
TCAACGTAGG CTGAACACCA GAGCGACAAG GAACCCTCCC AGAGGAGACT CACTAACTGA

3001 ACCCGTCAGC GGGGGTCTTT CACACATGCA GCATGTATCA AAATTAATTT GGTITTTTTT
TGGCAGTCC CCCCAGAAA GTGTGTACGT CGTACATAGT TTTAATTAAT CCAAAAAA

3061 CTTAAGTATT TACATTAAAT GGCCATAGTA CTTAAAGTTA CATTGGCTTC CTTGAAATAA
GAATTCATAA ATGTAATTTA CCGGTATCAT GAATTTCAAT GTAACCGAAG GAACTTTATT

3121 ACATGGAGTA TTCAGAATGT GTCATAAATA TTTCTAATTT TAAGATAGTA TCTCCATTGG
TGTACCTCAT AAGTCTTACA CAGTATTTAT AAAGATTAAA ATTCTATCAT AGAGGTAACC

3181 CTTTCTACTT TTTCTTTTAT TTTTTTTTGT CCTCTGTCTT CCATTTGTTG TTGTTGTTGT
GAAAGATGAA AAAGAAAATA AAAAAAACA GGAGACAGAA GGTAACAAC AACAACAACA

FIG. 17D

APPROVED	O.C.
BY	CLASS
	DATE/TIME

3241 TTGTTTGTTT GTTGTGTGGT TGGTGTGTTA ATTTTTTTTT AAAGATCCTA CACTATAGTT
AACAAACAAA CAAACAACCA ACCAACCAAT TAAAAAATAA TTTCTAGGAT GTGATATCAA

3301 CAAGCTAGAC TATTAGCTAC TCTGTAACCC AGGGTGACCT TGAAGTCATG GGTAGCCTGC
GTTTCGATCTG ATAATCGATG AGACATTGGG TCCCACTGGA ACTTCAGTAC CCATCGGACC

3361 TGTTTTAGCC TTCCACATC TAAGATTACA GGTATGAGCT ATCATTTTTG GTATATTGAT
ACAAAATCGG AAGGGTGTAG ATTCTAATGT CCATACTCGA TAGTAAAAAC CATATAACTA

3421 TGATTGATTG ATTGATGTGT GTGTGTGTGA TTGTGTTTGT GTGTGTGANT GTGWANATGT
ACTAACTAAC TAACTACACA CACACACACT AACACAAACA CACACACTNA CACWTNTACA

3481 GTGTATGGNT GTGTGTGAKT GTGTGTATGT ATGNYTGTGT GTGANTGYGT GTGTGTGANT
CACATACCNA CACACACTMA CACACATACA TACNRACACA CACTNACRCA CACACACTNA

3541 GTGCATGTGT GTGTGTGTGA CTGTGTCTAT GTGTATGACT GTGTGTGTGT GTGTGTGTGT
CACGTACACA CACACACACT GACACAGATA CACATACTGA CACACACACA CACACACACA

3601 GTGTGTGTGT GTGTGTGTGT GTGTGTTGTG AAAAAATATT CTATGGTAGT GAGAGCCAAC
CACACACACA CACACACACA CACACAACAC TTTTTTATAA GATACCATCA CTCTCGGTTG

3661 GCTCCGGCTC AGGTGTCAGG TTGGTTTTTG AGACAGAGTC TTCACTTAG CTTGGAATTC
CGAGGCCGAG TCCACAGTCC AACCAAAAAAC TCTGTCTCAG AAAGTGAATC GAACCTTAAG

3721 TTGAAGACGA AAGGGCCTCG TGATACGCCT ATTTTTATAG GTTAATGTCA TGATAATAAT
AACTTCTGCT TTCCCGGAGC ACTATGCGGA TAAAAATATC CAATTACAGT ACTATTATTA

3781 GGTTCCTTAG ACGTCAGGTG GCACTTTTCG GGGAAATGTG CGCGGAACCC CTATTTGTTT
CCAAAGAATC TGCAGTCCAC CGTGAAAAGC CCCTTTACAC GCGCCTTGGG GATAAACAAA

3841 ATTTTTCTAA ATACATTCAA ATATGTATCC GCTCATGAGA CAATAACCTT GATAAATGCT
TAAAAAGATT TATGTAAGTT TATACATAGG CGAGTACTCT GTTATTGGGA CTATTACGA

3901 TCAATAATAT TAAAAAGGA AGAGTATGAG TATTCAACAT TTCCGTGTG CCGTTATTCC
AGTTATTATA ACTTTTTCCT TCTCATACTC ATAAGTTGTA AAGGCACAGC GGAATAAGG

3961 CTTTTTTGCG GCATTTTGCC TTCCTGTTTT TGCTCACCCA GAAACGCTGG TGAAAGTAAA
GAAAAACCGC CGTAAAACGG AAGGACAAAA ACGAGTGGGT CTTTGGGACC ACTTTCATTT

4021 AGATGCTGAA GATCAGTTGG GTGCACGAGT GGGTTACATC GAACTGGATC TCAACAGCGG
TCTACGACTT CTAGTCAACC CACGTGCTCA CCCAATGTAG CTTGACCTAG AGTTGTGCGC

FIG. 17E

4081 TAAGATCCTT GAGAGTTTTT CCCCCGAAGA ACGTTTTCCA ATGATGAGCA CTTTTAAAGT
ATTCTAGGAA CTCTCAAAAG CGGGGCTTCT TGCAAAAGGT TACTACTCGT GAAAATTTCA

4141 TCTGCTATGT GGCGCGGTAT TATCCCGTGT TGACGCCGGG CAAGAGCAAC TCGGTCGCCG
AGACGATACA CCGCGCCATA ATAGGGCACA ACTGCGGGCC GTTCTCGTTG AGCCAGCGGC

4201 CATACTAT TCTCAGAATG ACTTGTTGA GTACTACCA GTCACAGAAA AGCATCTTAC
GTATGTGATA AGAGTCTTAC TGAACCACT CATGAGTGGT CAGTGTCTTT TCGTAGAATG

4261 GGATGGCATG ACAGTAAGAG AATTATGCAG TGCTGCCATA ACCATGAGTG ATAACACTGC
CCTACCGTAC TGTATTCTC TTAATACGTC ACGACGGTAT TGGTACTCAC TATTGTGACC

4321 GGCCAACTTA CTTCTGACAA CGATCGGAGG ACCGAAGGAG CTAACCGCTT TTTTGCACAA
CCGGTTGAAT GAAGACTGTT GCTAGCCTCC TGGCTTCCTC GATTGGCGAA AAAACGTGTT

4381 CATGGGGGAT CATGTAATC GCCTTGATCG TTGGGAACCG GAGCTGAATG AAGCCATACC
GTACCCCTA GTACATTGAG CGGAAGTAGC AACCCTTGGC CTCGACTTAC TTCGGTATGG

4441 AAACGACGAG CGTGACACCA CGATGCCTGC AGCAATGGCA ACAACGTTGC GCAAACCTATT
TTTGCTGCTC GCACTGTGGT GCTACGGACG TCGTTACCGT TGTGCAACG CGTTTGATAA

4501 AACTGGCGAA CTAATTACTC TAGCTTCCCG GCAACAATTA ATAGACTGGA TGGAGGCGGA
TTGACCGCTT GATGAATGAG ATCGAAGGGC CGTTGTTAAT TATCTGACCT ACCTCCGCTT

4561 TAAAGTTGCA GGACCACTTC TCGGCTCGGC CTTTCCGGCT GGCTGGTTTA TTGCTGATAA
ATTTCAACGT CCTGGTGAAG ACCGAGCCG GGAAGGCCGA CCGACCAAAT AACCCTATT

4621 ATCTGGAGCC GGTGAGCGTG GGTCTCGCG TATCATTGCA GCACTGGGGC CAGATGGTAA
TAGACCTCGG CCACTCGCAC CCAGAGCGCC ATAGTAACGT CGTGACCCCG GTCTACCATT

4681 GCCCTCCCGT ATCGTAGTTA TCTACACGAC GGGGAGTCAG GCAACTATGG ATGAACGAAA
CGGGAGGGCA TAGCATCAAT AGATGTGCTG CCCCTCAGTC CGTTGATACC TACTTGCTTT

4741 TAGACAGATC GCTGAGATAG GTCCCTCACT GATTAAGCAT TGGTAACTGT CAGACCAAGT
ATCTGTCTAG CGACTCTATC CACGGAGTGA CTAATTCGTA ACCATTGACA GTCTGGTTCA

FIG. 17F

4801 TTA_{CT}CA_{TAT} ATA_{CTTT}AGA TTGA_{TTT}AAA ACT_{TCAT}TTT TAAT_{TTA}AAA GGAT_{CTAG}GT
AAT_{GAGT}ATA TAT_{GAA}ATCT AACT_{AAAT}TT AGA_{AGT}AAAA ATTA_{AAAT}TTT CCT_{AGAT}CCA

4861 GAAG_{ATC}CTT TTT_{GATA}ATC TCAT_{GAC}CAA AAT_{CCCT}TAA CGT_{GAGT}TTT CGT_{TCCA}CTG
CTT_{CTAG}GAA AA_{ACTAT}TAG AGT_{ACTG}GTT TTAG_{GGA}ATT GCA_{CTCA}AAA GCA_{AGGT}GAC

4921 AGCG_{TCAG}AC CCC_{GTA}AAA AGAT_{CAA}AGG ATCT_{TCT}TGA GAT_{CCT}TTTT TTCT_{GCG}CGT
TCG_{CAGT}CTG GGC_{ATC}TTT TCT_{AGT}TTCC TAGA_{AGA}ACT CTAG_{GAAAA} AAG_{ACG}CGCA

4981 AAT_{CTG}CTGC TTG_{CAA}ACAA AAAA_{ACC}ACC GCT_{ACC}AGCG GTG_{GTT}TGTT TGCC_{GGAT}CA
TTAG_{ACG}ACG AAC_{GTT}TGTT TTTT_{TGT}GCG CGAT_{GGT}CCG CAC_{CAA}ACAA ACC_{GCCT}AGT

5041 AGAG_{CTAC}CA ACT_{CTTT}TC CGA_{AGGT}AAC TGG_{CTT}CAGC AGAG_{CGC}AGA TAC_{CAA}ATAC
TCT_{GAT}GGT TGAG_{AAAA}AAG GCT_{TCCA}TTC ACC_{GAGT}CG TCT_{GCGT}CT ATG_{GTTT}ATG

5101 TGT_{CCTT}CTA GTG_{TAG}CCGT AGT_{TAG}GCCA CCA_{CTT}CAAG AACT_{CTGT}AG CAC_{CGC}CTAC
ACAG_{GAA}GAT CAC_{ATC}GCGA TCA_{ATC}CGT GGT_{GAA}GTTT TTG_{AGAC}ATC GTG_{GCG}GATG

5161 ATAC_{CTCG}CT CTG_{CTA}ATCC TGT_{TACC}AGT GGCT_{GCT}GCC AGT_{GCG}GATA AGT_{CGT}GTCT
TAT_{GAG}CGA GAC_{GATT}AGG ACA_{ATG}GTC CCG_{ACG}ACGG TCAC_{CGT}TAT TCAG_{CAC}AGA

5221 TAC_{CGG}GTTG GACT_{CAAG}AC GAT_{AGT}TACC GGAT_{AAG}GCG CAG_{CGG}TCCG GCT_{GAA}CGGG
ATG_{GCCA}AC CTG_{AGT}TCTG CTAT_{CAAT}GCG CCT_{ATT}CCGC GTC_{GCC}AGCC CGA_{CTT}GCCC

5281 GGG_{TTCG}TGC ACAC_{AGCC}CA GCT_{TGG}AGCG AAC_{GAC}CTAC ACC_{GAA}CTGA GAT_{ACCT}ACA
CCCA_{AGCA}CG TGT_{GTC}GGGT CGA_{ACCT}CGC TTG_{CTG}GATG TGG_{CTT}GA_{CT} CTAT_{GGAT}GT

5341 GCG_{TGAG}CTA TGAG_{AAAG}CG CCAC_{GCT}TCC CGA_{AGG}GAGA AAG_{GCG}GACA GGT_{ATCC}GGT
CGC_{ACT}CGAT ACT_{CTTT}CGC GGT_{GCGA}AGG GCT_{TCC}CTCT TTCC_{GCT}GT CCAT_{AGGC}CA

5401 AAG_{CGG}CAGG GTC_{GGA}ACAG GAG_{AGC}GCAC GAG_{GGAG}CTT CCAG_{GGG}GAA ACG_{CCT}GGTA
TTC_{GCG}TCC CAG_{CCT}TGTC CTCT_{GCGT}G CTCC_{CTC}GAA GGT_{CCCC}CTT TCG_{GACC}AT

5461 TCT_{TTAT}AGT CCT_{GTC}GGGT TTC_{GCC}ACCT CTG_{ACTT}GAG CGT_{CGAT}TTT TGT_{GATG}CTC
AGAA_{ATAT}CA GGAC_{AGCC}CA AAG_{CGG}TGGA GACT_{GAA}CTC GCAG_{CTAA}AA AACT_{ACG}AG

5521 GTC_{AGG}GGGG CGG_{AGC}CTAT GGAA_{AAAC}GC CAG_{CAAC}GCG GCCT_{TTTT}TAC GGT_{TCCT}GCG
CAG_{TCCCC} GCCT_{CGGA}T CCT_{TTTT}TGCG GTC_{GTT}GCG CGG_{AAAA}ATG CCA_{AGG}ACCG

FIG. 17G

5581 CTTTGTGCTGG CCTTTTGCTC ACATGTTCTT TCCTGCGTTA TCCCCTGATT CTGTGGATAA
GAAAACGACC GGAACGAG TGTACAAGAA AGGACGCAAT AGGGGACTAA GACACCTATT

5641 CCGTATTACC GCCTTTGAGT GAGCTGATAC CGCTCGCCGC AGCCGAACGA CCGAGCGCAG
GGCATAATGG CGGAACTCA CTCGACTATG GCGAGCGGCG TCGGCTTGCT GGCTCGCGTC

5701 CGAGTCAGTG AGCGAGGAAG CGGAAGAGCG CCTGATGCGG TATTTTCTCC TTACGCATCT
GCTCAGTCAC TCGCTCCTTC GCCTTCTCGC GGACTACGCC ATAAAAGAGG AATGCGTAGA

5761 GTGCGGTATT TCACACCGCA TATGGTGAC TCTCAGTACA ATCTGCTCTG ATGCCGCATA
CACGCCATAA AGTGTGGCGT ATACCACGTG AGAGTCATGT TAGACGAGAC TACGGCGTAT

5821 GTTAAGCCAG TATACACTCC GCTATCGCTA CGTGAAGGG TCATGGCTGC GCGGCGACAC
CAATTCGGTC ATATGTGAGG CGATAGCGAT GCACTGACCC AGTACCGACG CCGGGCTGTG

5881 CCGCCAACAC CCGCTGACGC GCCCTGACGG GCTTGTCTGC TCCCGGCATC CGCTTACAGA
GGCGGTTGTG GCGGACTGCG CCGGACTGCC CGAACAGACG AGGGCCGTAG GCGAATGTCT

5941 CAAGCTGTGA CCGTCTCCGG GAGCTGCATG TGTCAGAGGT TTTACCGTC ATCACCAGAA
GTTTCGACACT GGCAGAGGCC CTCGACGTAC ACAGTCTCCA AAAGTGGCAG TAGTGGCTTT

6001 CGCGCGAGGC AGCTGCGGTA AAGCTCATCA GCGTGGTCGT GAAGCGATTG ACAGATGTCT
GCGCGCTCCG TCGACGCCAT TTCGAGTAGT CGCACCAGCA CTTGCTAAG TGTCTACAGA

6061 GCCTGTTTAT CCGCGTCCAG CTCGTTGAGT TTCTCCAGAA CCGTTAATGT CTGGCTTCTG
CGGACAAGTA GCGCGAGGTC GAGCAACTCA AAGAGGTCTT CGCAATTACA GACCGAAGAC

6121 ATAAAGCGGG CCATGTTAAG GGCGGTTTTT TCCTGTTTGG TCACTTGATG CCTCCGTGTA
TATTTGCCCC GGTACAATTG CCGCCAAAAA AGGACAAACC AGTGAAGTAC GGAGGCACAT

6181 AGGGGGAATT TCTGTTTATG GGGGTAATGA TACCGATGAA ACGAGAGAGG ATGCTCACGA
TCCCCCTTAA AGACAAGTAC CCCCATTACT ATGGCTACTT TGCTCTCTCC TACGAGTGCT

6241 TACGGGTTAC TGATGATGAA CATGCCCGGT TACTGGAACG TTGTGAGGGT AAACAAGTGG
ATGCCCAATG ACTACTACTT GTACGGGCCA ATGACCTTGC AACACTCCA TTTGTTGACC

6301 CCGTATGGAT GCGGCGGGAC CAGAGAAAAA TCACTCAGGG TCAATGCCAG CGCTTCGTTA
GCCATACCTA CGCGGCCCTG GTCTCTTTTT AGTGAGTCCC AGTTACGGTC GCGAAGCAAT

FIG. 17H

6361	ATACAGATGT TATGTCTACA	AGGTGTTCCA TCCACAAGGT	CAGGGTAGCC GTCCCATCGG	AGCAGCATCC TCGTCTAGG	TCCGATGCAG ACGCTACGTC	ATCCGGAACA TAGGCCTTGT
6421	TAATGGTGCA ATTACCACGT	GGGCGCTGAC CCCGCGACTG	TTCCGCGTTT AAGGCGCAAA	CCAGACTTTA GGTCTGAAAT	CGAAACACGG GCTTTGTGCC	AAACCGAAGA TTTGGCTTCT
6481	CCATTCATGT GGTAAGTACA	TGTTGCTCAG ACAACGAGTC	GTCGCAGACC CAGCGTCTGC	TTTTGCAGCA AAAACGTCGT	GCAGTCGCTT CGTCAGCGAA	CACGTTCCGT GTGCAAGCGA
6541	CGCGTATCGG GCGCATAGCC	TGATTCATTC ACTAAGTAAG	TGCTAACCAG ACGATTGGTC	TAAGGCAACC ATTCCGTTGG	CCGCCAGCCT GGCGGTCGGA	AGCCGGGTCC TCGGCCCAGG
6501	TCAACGACAG AGTTGCTGTC	GAGCACGATC CTCGTGCTAG	ATGCCGACCC TACGCGTGGG	GTGGCCAGGA CACCGGTCCT	CCCAACGCTG GGGTTGCGAC	CCCGAGATGC GGGCTCTACG
6561	GCCGCGTGCG CGGCGCACGC	GCTGCTGGAG CGACGACCTC	ATGGCGGACG TACCGCCTGC	CGATGGATAT GCTACCTATA	GTTCTGCCAA CAAGACGGTT	GGGTTGGTTT CCCAACCAAA
6721	GCGCATTAC CGCGTAAGTG	AGTTCTCCGC TCAAGAGGCG	AAGAATTGAT TTCTTAACTA	TGGCTCCAAT ACCGAGGTTA	TCTTGGAGTG AGAACCTCAC	GTGAATCCGT CACTTAGGCA
6781	TAGCGAGGTG ATCGTCCAC	CCGCCGGCTT GGCGGCCGAA	CCATTCAGGT GGTAAGTCCA	CGAGGTGGCC GCTCCACCGG	CGGCTCCATG GCCGAGGTAC	CACCGCGACG GTGGCGCTGC
6841	CAACGCGGGG GTTGCGCCCC	AGGCAGACAA TCCGTCTGTT	GGTATAGGGC CCATATCCCG	GGCGCCTACA CCGCGGATGT	ATCCATGCCA TAGGTACGGT	ACCCGTTCCA TGGGCAAGGT
6901	TGTGCTCGCC ACACGAGCGG	GAGGCGGCAT CTCCGCCGTA	AAATCGCCGT TTTAGCGGCA	GACGATCAGC CTGCTAGTCG	GGTCCAGTGA CCAGGTCACT	TCAAGTTAG AGCTTCAATC
6961	GCTGGTAAGA CGACCATTCT	GCCGCGAGCG CGGCGCTCGC	ATCCTTGAAG TAGGAAC TTC	CTGTCCCTGA GACAGGGACT	TGGTCGTCAT ACCAGCAGTA	CTACCTGCCT GATGGACGGC
7021	GGACAGCATG CCTGTCTGAC	GCCTGCAACG CGGACGTTGC	CGGGCATCCC GCCCCGTAGG	GATGCCGCCG CTACGGCGGC	GAAGCGAGAA CTTCGCTCTT	GAATCATAAT CTTAGTATTA
7081	GGGGAAGGCC CCCCTTCCGG	ATCCAGCCTC TAGGTCCGAG	GCGTCGCGAA CGCAGCGCTT	CGCCAGCAAG GCGGTCGTTT	ACGTAGCCCA TGCATCGGGT	GCGGTCGGC CGCGCAGCCG
7141	CGCCATGCCG GCGGTACGGC	GCGATAATGG CGCTATTACC	CCTGCTTCTC GGACGAAGAG	GCCGAAACGT CGGCTTTGCA	TTGGTGGCGG AACCACCGCC	GACCAGTGAC CTGGTCACTG

FIG. 17I

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7201 GAAGGCTTGA GCGAGGGCGT GCAAGATTCC GAATACCGCA AGCGACAGGC CGATCATCGT
CTTCCGAAC TCGTCCCGCA CGTTCTAAGG CTTATGGCGT TCGTGTCCG GCTAGTAGCA

7261 CGCGCTCCAG CGAAAGCGGT CCTCGCCGAA AATGACCCAG AGCGCTGCCG GCACCTGTCC
GCGCGAGGTC GCTTTCGCCA GGAGCGGCTT TTAGTGGGTC TCGCGACGGC CGTGGACAGG

7321 TACGAGTTGC ATGATAAAGA AGACAGTCAT AAGTGGGGCG ACGATAGTCA TGCCCCGGCG
ATGCTCAACG TACTATTTCT TCTGTCAGTA TTCACGGCGC TGCTATCAGT ACGGGGCGCG

7381 CCACCGGAAG GAGCTGACTG GGTGAAGGC TCTCAAGGGC ATCGGTCGAC GCTCTCCCTT
GGTGGCCTTC CTCGACTGAC CCAACTTCCG AGAGTTCCCG TAGCCAGCTG CGAGAGGGAA

7441 ATGCGACTCC TGCATTAGGA AGCAGCCCAG TAGTAGGTTG AGCCCGTTGA GCACCGCCGC
TACGCTGAGG ACCTAATCCT TCGTCGGGTC ATCATCCAAC TCCGGCAACT CGTGGCGGCG

7501 CGCAAGGAAT GGTGCATGCA AGGAGATGGC GCCAACAGT CCCCCGGCCA CGGGGCCTGC
GCGTTCCTTA CCACGTACGT TCCTCTACCG CGGGTTGTCA GGGGCGCGGT GCCCGGACG

7561 CACCATACCC ACGCCGAAAC AAGCGTCAT GAGCCCGAAG TGGCGAGCCC GATCTTCCCC
GTGGTATGGG TCGCGCTTTG TTCGCGAGTA CTCGGGCTTC ACCGCTCGGG CTAGAAGGGG

7621 ATCGGTGATG TCGGCGATAT AGGCGCCAGC AACCGCACCT GTGGCGCCCG TGATGCCGGC
TAGCCACTAC AGCCGCTATA TCCGCGGTCC TTGGCGTGGA CACCGCGGCC ACTACGGCCG

7681 CACGATGCGT CCGGCGTAGA GCGCCACAGG ACGGGTGTGG TCGCCATGAT CGCGTAGTCG
GTGCTACGCA GGCCGCATCT CGCGGTGTCC TGCCACACCC AGCGGTACTA GCGCATCAGC

7741 ATAGTGGCTC CAAGTAGCGA AGCGAGCAGG ACTGGGCGGC GGCCAAAGCG GTCGGACAGT
TATACCCGAG GTTCATCGCT TCGCTCGTCC TGACCCGCGC CCGGTTTCCG CAGCCTGTCA

7801 GCTCCGAGAA CGGGTGGCA TAGAAATTGC ATCAACGCAT ATAGCGCTAG CAGCAGCCA
CGAGGCTCTT GCCACGCGT ATCTTTAACG TAGTTGCGTA TATCGCGATC GTCGTGCGGT

7861 TAGTACTGG CGATGCTGTC GGAATGGACG ATATCCCGCA AGAGGCCCGG CAGTACCGGC
ATCACTGACC GCTACGACAG CCTTACCTGC TATAGGGCGT TCTCCGGGCC GTCATGGCCG

7921 ATAACCAAGC CTATGCCTAC AGCATCCAGG GTGACGGTGC CGAGGATGAC GATGAGCGCA
TATTGGTTCC GATACGGATG TCGTAGGTCC CACTGCCACG GCTCCTACTG CTAATCGGT

7981 TTGTTAGATT TCATACACGG TGCCTGACTG CGTTAGCAAT TTAAGTGTGA TAACTACCG
AACAATCTAA AGTATGTGCC ACGGACTGAC GCAATCGTTA AATTGACACT ATTTGATGGC

7041 CATTA
GTAAT

FIG. 17J